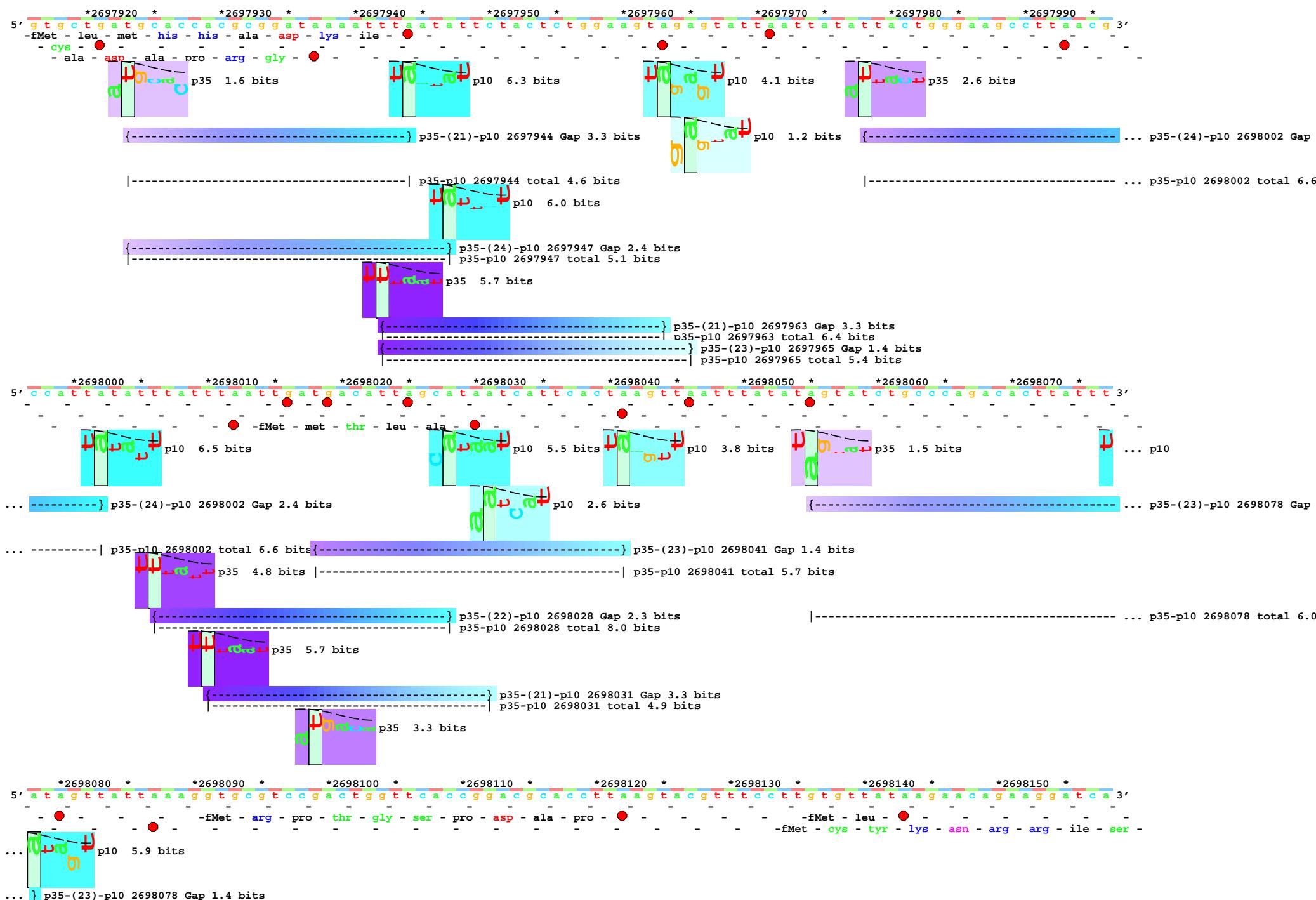
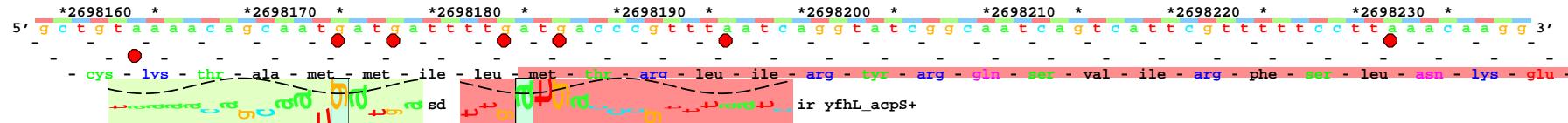


- 1 -  
piece 1, NC\_000913, yfhL\_acpS+, config: linear, direction: +, begin: 2697916, end: 2698659



... | p35-p10 2698078 total 6.0 bits



sd-(10)-ir 2698185 Gap 2.7 bits  
sd-ir 2698185 yfhL acsP+ total 6.5 bits

The diagram illustrates the gene structure of *yfhL\_acps+*. It shows the 5' to 3' directionality of the DNA sequence. Promoters are indicated by asterisks (\*), and transcription start sites are marked with red dots. The gene is transcribed from left to right. Two RNA secondary structures are shown: one in green labeled 'sd' (sigma factor binding site) and another in pink labeled 'ir yfhL\_acps+'. The pink structure represents an inverted repeat (IR) element.

[###] orf 3 codons  
sd-(12)-ir 2698367 Gap 4.0 bits  
sd-ir 2698367 yfhL\_acps+ total 5.7 bit

This figure shows the secondary structure of a tRNA precursor with various processing sites indicated by red dots. The mature tRNA regions are color-coded: p35 (purple), p10 (cyan), sd (light green), and ir (red). The precursor sequence is shown above the structure, with codons separated by hyphens. Processing sites are marked with red dots at positions 2698410, 2698420, 2698430, 2698440, 2698450, 2698460, 2698470, and 2698480. The p35 region is labeled as 3.5 bits, p10 as 5.2 bits, and sd as 4.5 bits. A gap of 4.0 bits is indicated between p35 and p10. The mature tRNA regions are labeled as p35-(25)-p10, sd, and ir.

|-----| p35-p10 2698439 total 4.7 bits |-----| p35 5.3 bits |-----| ... sd-ir 2698484 yfhL\_acpS+

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```
{-----} p35-(25)-p10 2698481 Gap 4.0 bits  
|-----| b35-p10 2698481 total 4.5 bits  
{-----} ... p35-(24)-p10 2698484 Gap 7.0 bits  
|-----| b35-p10 2698484 total 7.5 bits
```

5' \* \*2698490 \* \*2698500 \* \*2698510 \* \*2698520 \* \*2698530 \* \*2698540 \* \*2698550 \* \*2698560 3'  
 - fMet - leu - lys - ile - ala - ser - cys - leu - - fMet - leu - arg - val - his - ala - ala - gln -  
 - met - leu - lys - ile - ala - ser - cys - leu - - fMet - gln - val - ala - cys - lys - arg - - fMet - leu - his - lys -  
 ... ir yfhL\_acpS+ [###> orf 9 codons p10 5.8 bits p35 3.8 bits ...

... --> sd-(12)-ir 2698484 Gap 4.0 bits



...  p10 3.7 bits